

Fig. 1

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Molecule:

Sequence Data

Description:

chr15ed2.seq,

37544 bps DNA

1	ggatcagctg	gtttgccacc	gggatcccca	cgttgatgc	cctgttagcg	gaggaattct
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181	cgttgcaagg	ccaaccgtta	ccagctgagg	tctatggcga	cgatggcacc	gcgacggcgg
241	gtatccccta	taccgttatg	gacagtcggc	cccaggttcg	gcttctgacg	ggtttaccgg
301	gtaactcacc	gacagtcctg	ccgagtggtg	ttgaacagag	aacctggcag	tacgaaccgga
361	tcgccgatga	tccgcaatgc	catcagcagg	tggctgctga	cagtgaaccg	tacgggtttc
421	cacgggagac	cgtcgacatt	gcttatccgc	gccgccctaa	gcctgcgggtg	tcaccttacc
481	cggatacgct	gccggcgacg	ttattcgaca	gcagctatga	tgagcagcaa	tgagcaattgc
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721	tgacctatct	agggcatcaa	cgcgtggcct	ataccgggaa	gacccggaac	gaagaaaaac
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1981	cgctctctta	attaacacga	taacgttaaa	taatcacacc	ttcctgcoag	gtacggggga
2041	aggttancta	ctctatcaag	gaaagggttt	atgactgtaa	acagaggcga	taacctgcat
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2401	ctggccatca	atgcaaccca	agtcactcgt	acgtggcaat	atgagaatga	cactttacc
2461	ggacgcccgc	tcagtatcac	agaacagcct	gctggcgaag	caggccgtat	cacagagcgt
2521	tttgtcttgg	cagggaacag	tcaggcggag	aagaaacagca	acctggccgg	acagtgcgtg
2581	cgtaactatg	acaccccggt	actgaaccag	acggacagta	ttgcgcttaa	cggcataccg
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2761	gccacccggc	cggtactgac	caccacccgat	gcggccggta	acctgcagcg	tgtggcgtat
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3061	gaccgggttg	ggaaacgtgg	gaaagtgcag	aacgatgcgg	agggttaccg	cttctggcgc
3121	aaccaaaaag	tggctgccgg	gaacacctat	gtctatgaca	gcctgtatca	gctggtcagt
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3601	cagcaga gg	ctaacagctc	gcaggcgag	agcaecgtgt	acctgccagg	gctggagcgg
3661	caacaccaca	taaatggcac	gacgggtgaa	gaggtgctac	acgttatcac	gataggcgag
3721	gcgggcccgtg	cgcaggtgcg	ggtactgcac	tggaagaaac	gaaagccggg	tgccatcagt
3781	aacaaccaga	tgcgctacag	ctatgataac	cttatcggca	gcagcgggtc	ggaggtggac
3841	ggtgacggac	aaattatcag	tatggaagaa	tactacccgt	acggggggac	tgcggtgtgg
3901	acggcgagga	gtcagacaga	ggctgattac	aagactgtgc	gttactcagg	caaggagcgg

Fig. 2

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chr1m5ed2.seq

3961	gatgcaacgg	ggctgtatta	ttacggctac	cggtattacc	agccgtgggc	ggggagctgg
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4141	ccattttattc	atgaggacag	gatttttctg	ctggcaagcg	cgaatgttta	cagaacggaa
4201	cataataaat	ctgacatcat	tgcggttgta	gaagataaag	cattagatag	taaactattc
4261	accaatagta	tcgagcagtt	tttcaaaaaa	cctaaaggaa	aagcaatcct	gaaaggatcc
4321	cctgatatta	aagaaaggct	actcaataat	atagtacatg	acctgagcaa	tatgcaggta
4381	ggagatcagc	tgtatgtaaa	cgctcatggt	cattctgcga	aaccattttt	ttactccgat
4441	tcgggatatt	caaaatctat	catggaaacg	ctccaaagag	gggttaacta	tgtagctaaa
4501	gatttagtaa	ataagtttaa	attaccagaa	aatgcaacaa	tcaagataag	tacgtgtcat
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4861	gaaaacattt	ttgggtttaa	ggtaaaaaga	agctctgacg	gaacaaactt	taccggcagc
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6301	tgagtacata	gtccatcagg	cgcggtcgct	cccggaaggt	tcttcatcga	gnaattcgcc
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6601	cacattagca	catataatgg	tttgtggeta	tttattacac	agggggctag	atatcgtttt
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7081	taccaagggc	ggtattcctg	ccgcctgagg	gtttctgctg	attatccctt	tctgcctttt
7141	cccgcattct	gttagaatgc	ccacttctta	attgtttcca	aaactaatgt	tgttatgtca
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7381	ggtagccaaa	tttatgatct	cggttgctcc	cttggggcgg	caactctgtc	catccgcgcg
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8221	aaggcccttt	ttcattgtat	gacgttgaaa	ttgataccga	atggcgctct	gactggaaat

Fig. 2(i)

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chr1m5ed2.seq

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8581	aactgaaaaa	tcaactgggtg	tctgatgggtg	agttagtgtc	ggaaagttaa	gtgattgagg
8641	gtgatgaaaa	tcagtgcctc	attccgggtg	aacgctatgc	acaaatgcgg	aatgtctact
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8761	gaattcgtcga	tcattgcggct	acaacacctg	atgaacagcg	ccggacagaa	tggatgaaga
8821	ccgaatcact	ggtagatttc	cttgacctat	cagatcacag	taaaacaatt	gaaggctacc
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12061	gcgcggtgaa	atatgcagat	ctttccaaaga	gccgtactac	agactatgtt	ttcgactggg
12121	ataatatgct	ggcctttgaa	ggcaacacgg	caacttatat	gcaatacggc	tacacgcggc
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12361	gtctgttctc	tggtttctat	gagcactgcc	ctatcctgaa	tgccgatagc	gaagaaactgc
12421	gccagagccg	cctgaagttg	gctctgtga	cagcgnaaac	ttgaagcaa	ggctctgata
12481	ctctgggtat	tcagactgta	gaacgtatgt	ataatcttc	tacaagctg	aaaactggcg
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Fig. 2(ii)

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12721  aaaaaacotg  atttttattc  actatttaata  attaatgata  atttctattt  taattaacct
12781  tgttataaaa  aatagtattt  taaaaaaaca  ttttacatta  tataaaatat  atcaatcgac
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12901  tattatattt  acatcacgtt  agatcaaaat  aacaaattttt  tagtccgtta  cccagattca
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16741  aaataacgct  tccgtatggc  accatcgtaa  atcgcaata  tgcacccctt  agtactgata
16801  cgctgataac  agatattcga  gtgaatggaa  ttctctggg  acagcaaacg  tttgacgggt
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Fig. 2(iii)

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17161	tgagttacct	atggacactt	aggggtctgg	agaacggtta	cactgatctg	actggaacaa
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17281	agactactct	aaattacgat	gacctgaatc	gccatattgg	tagtcaagta	acagatttag
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17401	ggaaattgtg	tgatagctca	ggccatacgt	tagatatcca	gcagagctgg	ctgaaaaaac
17461	agcaattagc	aaatagaata	gtgaaactga	atggagttat	gcagcgtaca	gaacagttact
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17641	ccgcctgtca	caccacattc	gcagatggga	cagaagacca	tgctaccttc	aaatttgcca
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19141	tcttataaaag	tacttaccat	tcgtatgac	cccggaagat	catactggga	agaaacagaa
19201	aaacagtgaa	accaaanaact	tttcgcccct	ataaaccttg	caaaattcaa	aggtactggc
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Fig. 2(iv)

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Fig. 2(v)

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Fig. 2(vi)

SUBSTITUTE SHEET (RULE 26)

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Fig. 2(vii)

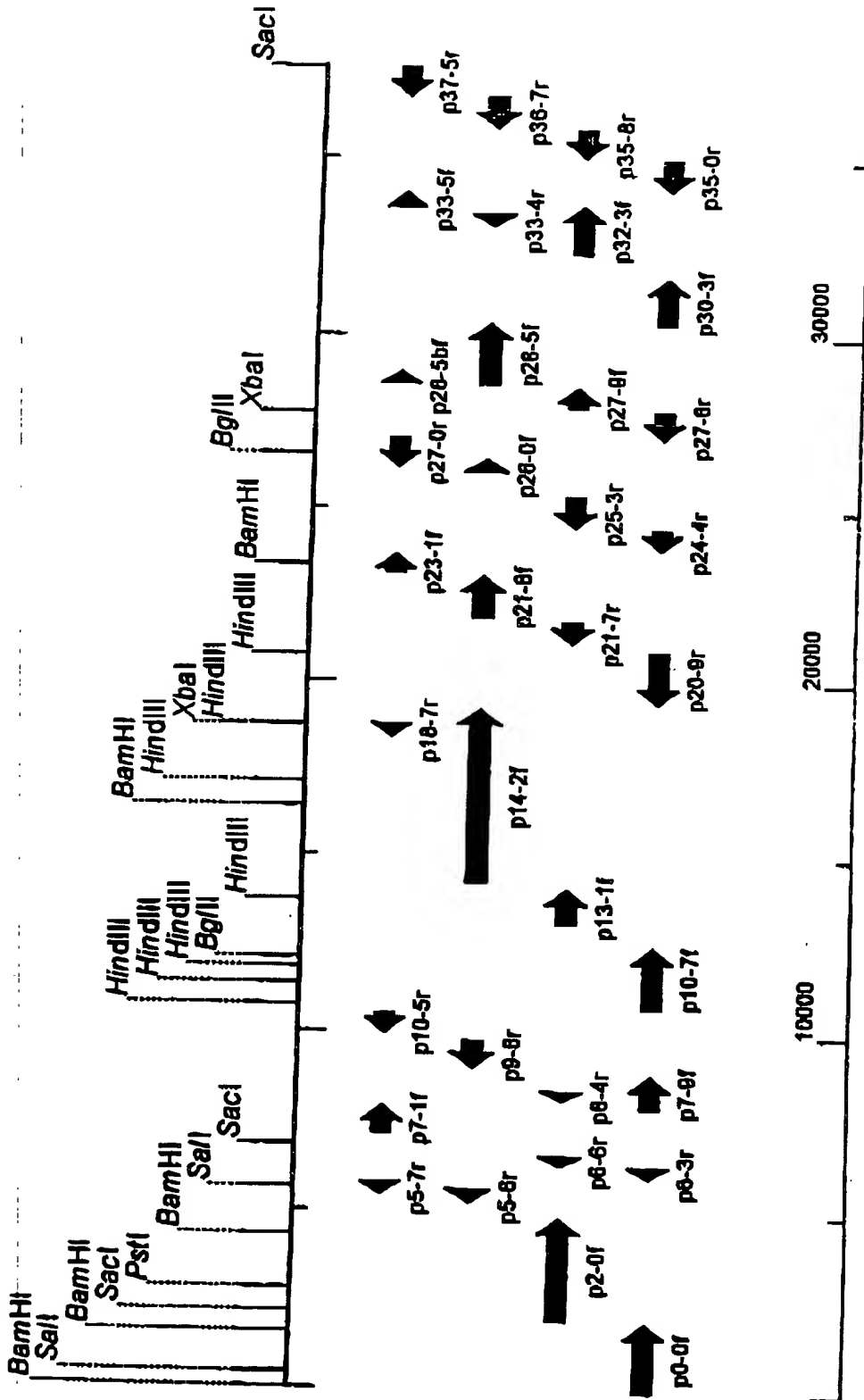
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Fig. 2(viii)

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chrom5ed2.seq (37544 bps) Fig. 3

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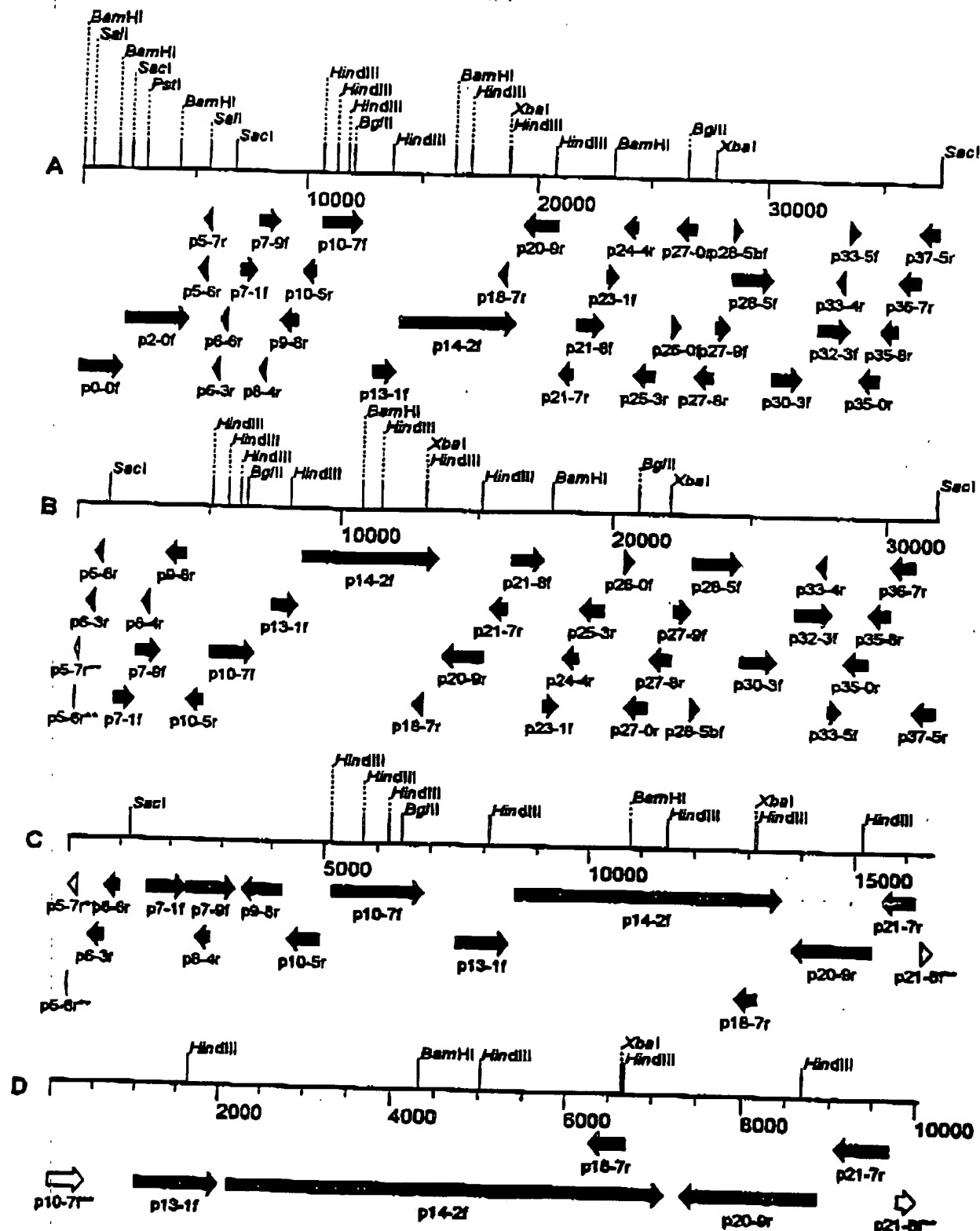


Fig. 4

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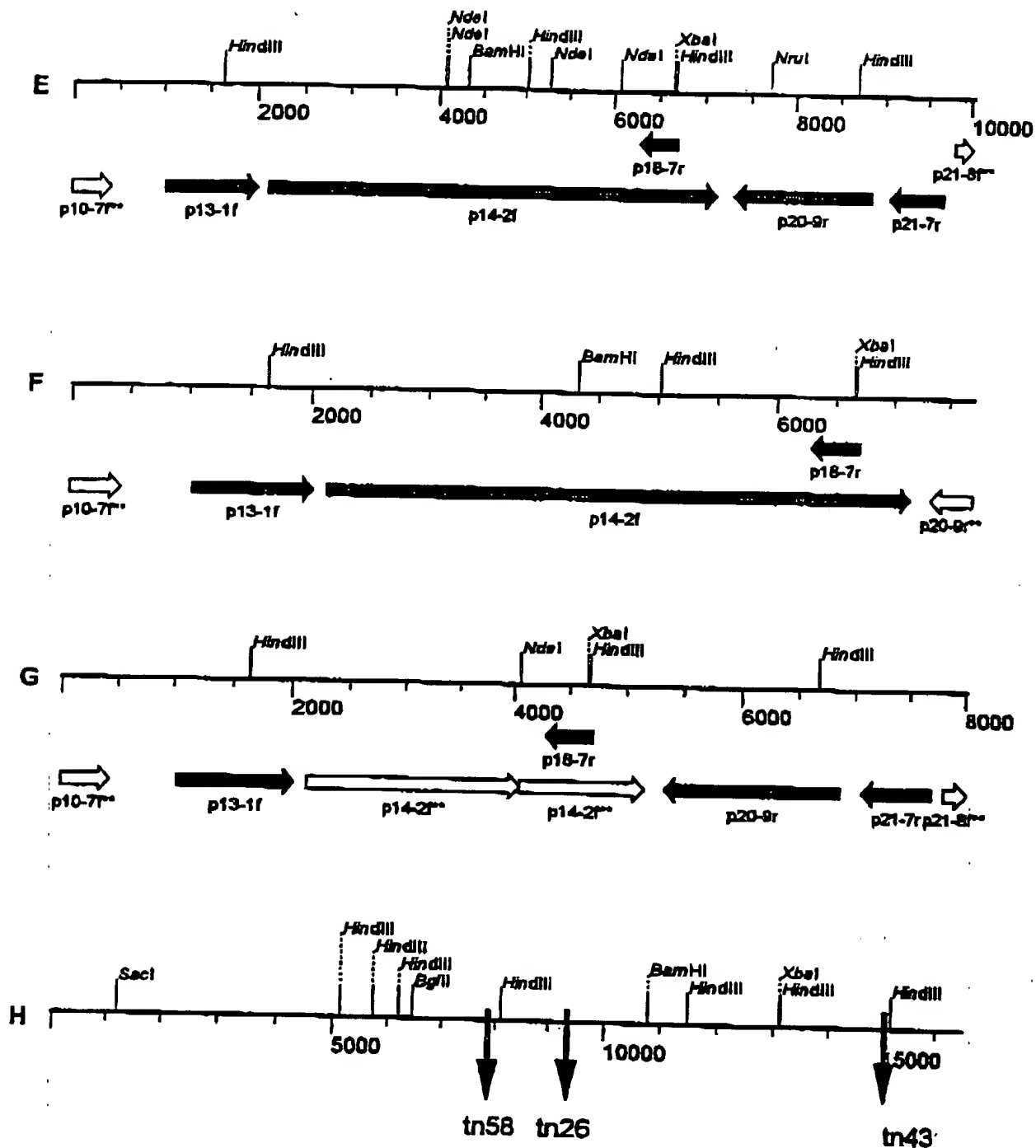


Fig. 4(cont'd)

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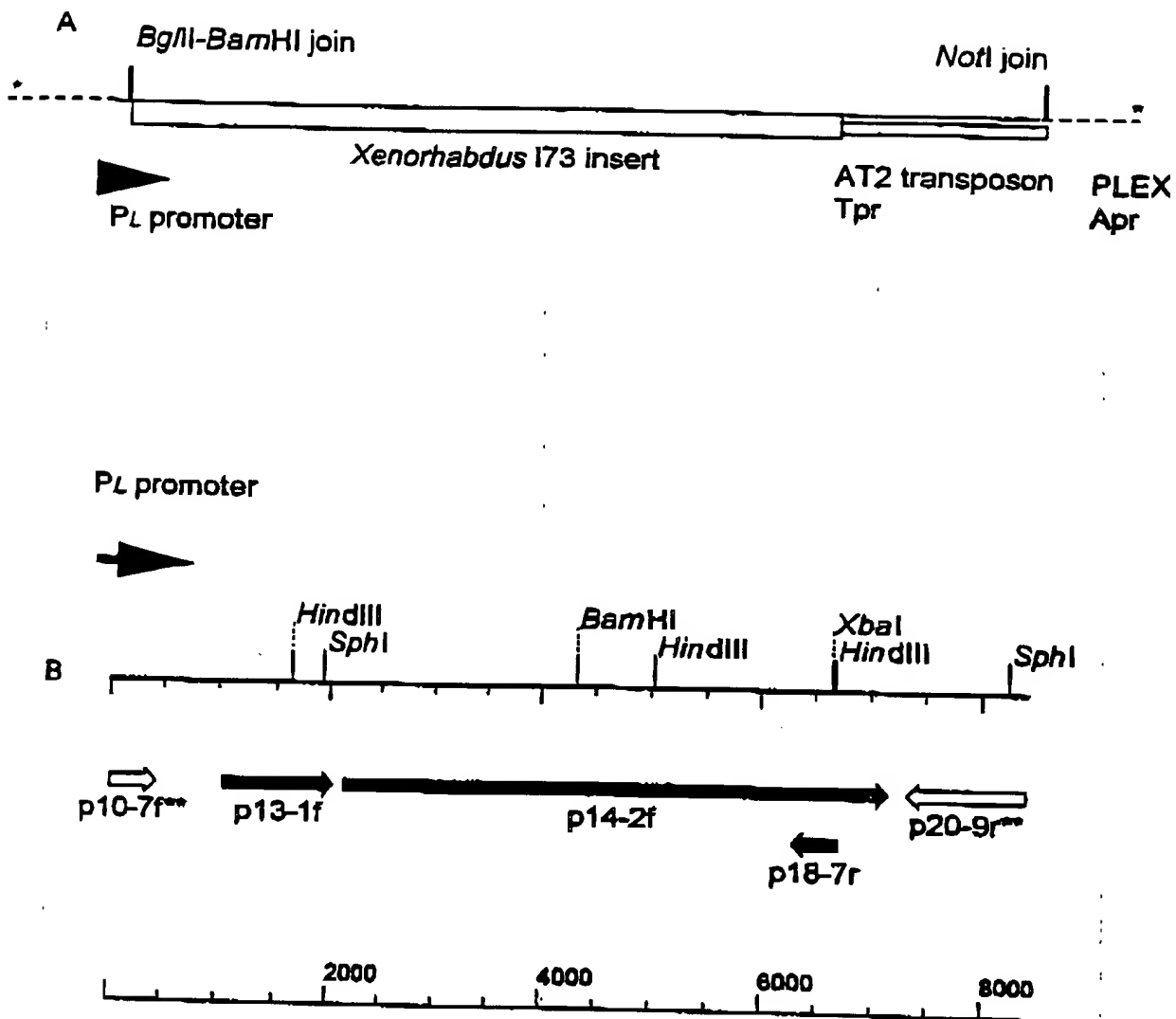


Fig. 5